SEQUENCE LISTING

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<120> COMPOUNDS AND METHODS FOR MODULATING ACTIVATION OF NF-kB

<130> 860098.427

<140> US
<141> 1998-12-10

<160> 30

<170> PatentIn Ver. 2.0

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Gly Leu Asp Ser Met Lys Asp Glu Glu Tyr Glu Gln Met Val Lys Glu
        35                  40                  45

Leu Gln Glu Ile Arg Leu Glu Pro Gln Glu Val Pro Arg Gly Ser Glu
    50                  55                  60

Pro Trp Lys Gln Gln Leu Thr Glu Asp Gly Asp Ser Phe Leu His Leu
65                  70                  75                  80

Ala Ile Ile His Glu Glu Lys Ala Leu Thr Met Glu Val Ile Arg Gln
                85                  90                  95

Val Lys Gly Asp Leu Ala Phe Leu Asn Phe Gln Asn Asn Leu Gln Gln
            100                 105                 110

Thr Pro Leu His Leu Ala Val Ile Thr Asn Gln Pro Glu Ile Ala Glu
        115                 120                 125

Ala Leu Leu Gly Ala Gly Cys Asp Pro Glu Leu Arg Asp Phe Arg Gly
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Asn Thr Pro Leu His Leu Ala Cys Glu Gln Gly Cys Leu Ala Ser Val
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Gly Val Leu Thr Gln Ser Cys Thr Thr Pro His Leu His Ser Ile Leu
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Lys Ala Thr Asn Tyr Asn Gly His Thr Cys Leu His Leu Ala Ser Ile
            180                 185                 190

His Gly Tyr Leu Gly Ile Val Glu Leu Leu Val Ser Leu Gly Ala Asp
        195                 200                 205

Val Asn Ala Gln Glu Pro Cys Asn Gly Arg Thr Ala Leu His Leu Ala
    210                 215                 220

Val Asp Leu Gln Asn Pro Asp Leu Val Ser Leu Leu Leu Lys Cys Gly
225                 230                 235                 240

Ala Asp Val Asn Arg Val Thr Tyr Gln Gly Tyr Ser Pro Tyr Gln Leu
                245                 250                 255

Thr Trp Gly Arg Pro Ser Thr Arg Ile Gln Gln Gln Leu Gly Gln Leu
            260                 265                 270

Thr Leu Glu Asn Leu Gln Met Leu Pro Glu Ser Glu Asp Glu Glu Ser
        275                 280                 285

Tyr Asp Thr Glu Ser Glu Phe Thr Glu Phe Thr Glu Asp Glu Leu Pro
    290                 295                 300

Tyr Asp Asp Cys Val Phe Gly Gly Gln Arg Leu Thr Leu
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Gly Pro Gly Leu Gly Ala Glu Leu Gly Pro Glu Leu Ser Trp Ala Pro
        35                  40                  45

Leu Val Phe Gly Tyr Val Thr Glu Asp Gly Asp Thr Ala Leu His Leu
    50                  55                  60

Ala Val Ile His Gln His Glu Pro Phe Leu Asp Phe Leu Leu Gly Phe
65                  70                  75                  80

Ser Ala Gly His Glu Tyr Leu Asp Leu Gln Asn Asp Leu Gly Gln Thr
                85                  90                  95

Ala Leu His Leu Ala Ala Ile Leu Gly Glu Ala Ser Thr Val Glu Lys
            100                 105                 110

Leu Tyr Ala Ala Gly Ala Gly Val Leu Val Ala Glu Arg Gly Gly His
        115                 120                 125

Thr Ala Leu His Leu Ala Cys Arg Val Arg Ala His Thr Cys Ala Cys
    130                 135                 140

Val Leu Leu Gln Pro Arg Pro Ser His Pro Arg Asp Ala Ser Asp Thr
145                 150                 155                 160

Tyr Leu Thr Gln Ser Gln Asp Cys Thr Pro Asp Thr Ser His Ala Pro
                165                 170                 175

Ala Ala Val Asp Ser Gln Pro Asn Pro Glu Asn Glu Glu Glu Pro Arg
            180                 185                 190
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Asp Glu Asp Trp Arg Leu Gln Leu Glu Ala Glu Asn Tyr Asp Gly His
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Thr Pro Leu His Val Ala Val Ile His Lys Asp Ala Glu Met Val Arg
    210                 215                 220

Leu Leu Arg Asp Ala Gly Ala Asp Leu Asn Lys Pro Glu Pro Thr Cys
225                 230                 235                 240

Gly Arg Thr Pro Leu His Leu Ala Val Glu Ala Gln Ala Ala Ser Val
                245                 250                 255

Leu Glu Leu Leu Leu Lys Ala Gly Ala Asp Pro Thr Ala Arg Met Tyr
            260                 265                 270

Gly Gly Arg Thr Pro Leu Gly Ser Ala Leu Leu Arg Pro Asn Pro Ile
        275                 280                 285

Leu Ala Arg Leu Leu Arg Ala His Gly Ala Pro Glu Pro Glu Asp Glu
    290                 295                 300

Asp Asp Lys Leu Ser Pro Cys Ser Ser Ser Gly Ser Asp Ser Asp Ser
305                 310                 315                 320

Asp Asn Arg Asp Glu Gly Asp Glu Tyr Asp Asp Ile Val Val His Ser
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Lys Asp Glu Glu
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Lys Arg Arg

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Ala Met Lys Asp Glu Glu
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Pro Asp Tyr Ala Met Tyr Pro Tyr Asp Val Pro Asp Tyr Ala Met Phe
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Gln Ala Ala Glu Arg Pro Gln Glu Trp Ala Met Glu Gly Pro Arg Asp
35 40 45

Gly Leu Lys Lys Glu Arg Leu Leu Asp Asp Arg His Asp Ser Gly Leu
50 55 60

Asp Ser Met Lys Asp Glu Glu Tyr Glu Gln Met Val Lys Glu Leu Gln
65 70 75 80

Glu Ile Arg Leu Glu Pro Gln Glu Val Pro Arg Gly Ser Glu Pro Trp
85 90 95

Lys Gln Gln Leu Thr Glu Asp Gly Asp Ser Phe Leu His Leu Ala Ile
100 105 110

Ile His Glu Glu Lys Ala Leu Thr Met Glu Val Ile Arg Gln Val Lys
115 120 125

Gly Asp Leu Ala Phe Leu Asn Phe Gln Asn Asn Leu Gln Gln Thr Pro
130 135 140

Leu His Leu Ala Val Ile Thr Asn Gln Pro Glu Ile Ala Glu Ala Leu
145 150 155 160

Leu Gly Ala Gly Cys Asp Pro Glu Leu Arg Asp Phe Arg Gly Asn Thr
165 170 175

Pro Leu His Leu Ala Cys Glu Gln Gly Cys Leu Ala Ser Val Gly Val
180 185 190

Leu Thr Gln Ser Cys Thr Thr Pro His Leu His Ser Ile Leu Lys Ala
195 200 205

Thr Asn Tyr Asn Gly His Thr Cys Leu His Leu Ala Ser Ile His Gly
210 215 220

Tyr Leu Gly Ile Val Glu Leu Leu Val Ser Leu Gly Ala Asp Val Asn
225 230 235 240

Ala Gln Glu Pro Cys Asn Gly Arg Thr Ala Leu His Leu Ala Val Asp
245 250 255

Leu Gln Asn Pro Asp Leu Val Ser Leu Leu Leu Lys Cys Gly Ala Asp
260 265 270

Val Asn Arg Val Thr Tyr Gln Gly Tyr Ser Pro Tyr Gln Leu Thr Trp
275 280 285

Gly Arg Pro Ser Thr Arg Ile Gln Gln Gln Leu Gly Gln Leu Thr Leu
290 295 300

Glu Asn Leu Gln Met Leu Pro Glu Ser Glu Asp Glu Glu Ser Tyr Asp
305 310 315 320

Thr Glu Ser Glu Phe Thr Glu Phe Thr Glu Asp Glu Leu Pro Tyr Asp
325 330 335

Asp Cys Val Phe Gly Gly Gln Arg Leu Thr Leu
340 345

<210> 13
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<213> Homo sapiens

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Pro Asp Tyr Ala Met Tyr Pro Tyr Asp Val Pro Asp Tyr Ala Met Phe
            20                  25                  30

Gln Ala Ala Glu Arg Pro Gln Glu Trp Ala Met Glu Gly Pro Arg Asp
        35                  40                  45

Gly Leu Lys Lys Glu Arg Leu Leu Asp Asp Arg His Asp Ala Gly Leu
    50                  55                  60

Asp Ala Met Lys Asp Glu Glu Tyr Glu Gln Met Val Lys Glu Leu Gln
65                  70                  75                  80

Glu Ile Arg Leu Glu Pro Gln Glu Val Pro Arg Gly Ser Glu Pro Trp
                85                  90                  95

Lys Gln Gln Leu Thr Glu Asp Gly Asp Ser Phe Leu His Leu Ala Ile
            100                 105                 110

Ile His Glu Glu Lys Ala Leu Thr Met Glu Val Ile Arg Gln Val Lys
        115                 120                 125

Gly Asp Leu Ala Phe Leu Asn Phe Gln Asn Asn Leu Gln Gln Thr Pro
    130                 135                 140

Leu His Leu Ala Val Ile Thr Asn Gln Pro Glu Ile Ala Glu Ala Leu
145                 150                 155                 160

Leu Gly Ala Gly Cys Asp Pro Glu Leu Arg Asp Phe Arg Gly Asn Thr
                165                 170                 175

Pro Leu His Leu Ala Cys Glu Gln Gly Cys Leu Ala Ser Val Gly Val
            180                 185                 190

Leu Thr Gln Ser Cys Thr Thr Pro His Leu His Ser Ile Leu Lys Ala
        195                 200                 205

Thr Asn Tyr Asn Gly His Thr Cys Leu His Leu Ala Ser Ile His Gly
    210                 215                 220

Tyr Leu Gly Ile Val Glu Leu Leu Val Ser Leu Gly Ala Asp Val Asn
225                 230                 235                 240

Ala Gln Glu Pro Cys Asn Gly Arg Thr Ala Leu His Leu Ala Val Asp
                245                 250                 255

Leu Gln Asn Pro Asp Leu Val Ser Leu Leu Leu Lys Cys Gly Ala Asp
            260                 265                 270

Val Asn Arg Val Thr Tyr Gln Gly Tyr Ser Pro Tyr Gln Leu Thr Trp
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275 280 285

Gly Arg Pro Ser Thr Arg Ile Gln Gln Leu Gly Gln Leu Thr Leu
290 295 300

Glu Asn Leu Gln Met Leu Pro Glu Ser Glu Asp Glu Glu Ser Tyr Asp
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Thr Glu Ser Glu Phe Thr Glu Phe Thr Glu Asp Glu Leu Pro Tyr Asp
325 330 335

Asp Cys Val Phe Gly Gly Gln Arg Leu Thr Leu
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35 40 45

Ser Gly Leu Gly Ser Leu Gly Pro Asp Ala Ala Ala Pro Gly Gly Pro
50 55 60

Gly Leu Gly Ala Glu Leu Gly Pro Glu Leu Ser Trp Ala Pro Leu Val
65 70 75 80

Phe Gly Tyr Val Thr Glu Asp Gly Asp Thr Ala Leu His Leu Ala Val
85 90 95

Ile His Gln His Glu Pro Phe Leu Asp Phe Leu Leu Gly Phe Ser Ala
100 105 110

Gly His Glu Tyr Leu Asp Leu Gln Asn Asp Leu Gly Gln Thr Ala Leu
115 120 125

His Leu Ala Ala Ile Leu Gly Glu Ala Ser Thr Val Glu Lys Leu Tyr
130 135 140

Ala Ala Gly Ala Gly Val Leu Val Ala Glu Arg Gly Gly His Thr Ala
145 150 155 160

Leu His Leu Ala Cys Arg Val Arg Ala His Thr Cys Ala Cys Val Leu
165 170 175

Leu Gln Pro Arg Pro Ser His Pro Arg Asp Ala Ser Asp Thr Tyr Leu

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          180                 185                 190

Thr Gln Ser Gln Asp Cys Thr Pro Asp Thr Ser His Ala Pro Ala Ala
        195                 200                 205

Val Asp Ser Gln Pro Asn Pro Glu Asn Glu Glu Glu Pro Arg Asp Glu
    210                 215                 220

Asp Trp Arg Leu Gln Leu Glu Ala Glu Asn Tyr Asp Gly His Thr Pro
225                 230                 235                 240

Leu His Val Ala Val Ile His Lys Asp Ala Glu Met Val Arg Leu Leu
                245                 250                 255

Arg Asp Ala Gly Ala Asp Leu Asn Lys Pro Glu Pro Thr Cys Gly Arg
            260                 265                 270

Thr Pro Leu His Leu Ala Val Glu Ala Gln Ala Ala Ser Val Leu Glu
        275                 280                 285

Leu Leu Leu Lys Ala Gly Ala Asp Pro Thr Ala Arg Met Tyr Gly Gly
    290                 295                 300

Arg Thr Pro Leu Gly Ser Ala Leu Leu Arg Pro Asn Pro Ile Leu Ala
305                 310                 315                 320

Arg Leu Leu Arg Ala His Gly Ala Pro Glu Pro Glu Asp Glu Asp Asp
                325                 330                 335

Lys Leu Ser Pro Cys Ser Ser Ser Gly Ser Asp Ser Asp Ser Asp Asn
            340                 345                 350

Arg Asp Glu Gly Asp Glu Tyr Asp Asp Ile Val Val His Ser Gly Arg
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Ser Gln Asn Arg Gln Pro Pro Ser Pro Ala Ser Lys Pro Leu Pro Asp
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Asp Pro Asn Pro Ala
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taaattgtac agatattatt catgacagct ttgtactgtg gaatgtgctt aataaaaaac 4080
aaaaaagttt gacttttgtc cagtaaattg ctaagtaatg tcaataaatc gagtatgggt 4140
attatgcagt gcacctaatc tggcttcatg caattgttac ttcagctact gattcaaagc 4200
caatactctt aataaagtgt tgcaatactc                                  4230
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<210> 16
<211> 542
<212> PRT
<213> Homo sapiens

<400> 16

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Met Glu Pro Asp Ser Val Ile Glu Asp Lys Thr Ile Glu Leu Met Cys
  1               5                  10                  15

Ser Val Pro Arg Ser Leu Trp Leu Gly Cys Ala Asn Leu Val Glu Ser
             20                  25                  30

Met Cys Ala Leu Ser Cys Leu Gln Ser Met Pro Ser Val Arg Cys Leu
         35                  40                  45

Gln Ile Ser Asn Gly Thr Ser Ser Val Ile Val Ser Arg Lys Arg Pro
     50                  55                  60

Ser Glu Gly Asn Tyr Gln Lys Glu Lys Asp Leu Cys Ile Lys Tyr Phe
 65                  70                  75                  80

Asp Gln Trp Ser Glu Ser Asp Gln Val Glu Phe Val Glu His Leu Ile
                 85                  90                  95

Ser Arg Met Cys His Tyr Gln His Gly His Ile Asn Ser Tyr Leu Lys
            100                 105                 110

Pro Met Leu Gln Arg Asp Phe Ile Thr Ala Leu Pro Glu Gln Gly Leu
        115                 120                 125

Asp His Ile Ala Glu Asn Ile Leu Ser Tyr Leu Asp Ala Arg Ser Leu
    130                 135                 140

Cys Ala Ala Glu Leu Val Cys Lys Glu Trp Gln Arg Val Ile Ser Glu
145                 150                 155                 160

Gly Met Leu Trp Lys Lys Leu Ile Glu Arg Met Val Arg Thr Asp Pro
                165                 170                 175

Leu Trp Lys Gly Leu Ser Glu Arg Arg Gly Trp Asp Gln Tyr Leu Phe
            180                 185                 190

Lys Asn Arg Pro Thr Asp Gly Pro Pro Asn Ser Phe Tyr Arg Ser Leu
        195                 200                 205
```

```
Tyr Pro Lys Ile Ile Gln Asp Ile Glu Thr Ile Glu Ser Asn Trp Arg
    210                 215                 220

Cys Gly Arg His Asn Leu Gln Arg Ile Gln Cys Arg Ser Glu Asn Ser
225                 230                 235                 240

Lys Gly Val Tyr Cys Leu Gln Tyr Asp Asp Glu Lys Ile Ile Ser Gly
                245                 250                 255

Leu Arg Asp Asn Ser Ile Lys Ile Trp Asp Lys Thr Ser Leu Glu Cys
            260                 265                 270

Leu Lys Val Leu Thr Gly His Thr Gly Ser Val Leu Cys Leu Gln Tyr
        275                 280                 285

Asp Glu Arg Val Ile Val Thr Gly Ser Ser Asp Ser Thr Val Arg Val
    290                 295                 300

Trp Asp Val Asn Thr Gly Glu Val Leu Asn Thr Leu Ile His His Asn
305                 310                 315                 320

Glu Ala Val Leu His Leu Arg Phe Ser Asn Gly Leu Met Val Thr Cys
                325                 330                 335

Ser Lys Asp Arg Ser Ile Ala Val Trp Asp Met Ala Ser Ala Thr Asp
            340                 345                 350

Ile Thr Leu Arg Arg Val Leu Val Gly His Arg Ala Ala Val Asn Val
        355                 360                 365

Val Asp Phe Asp Asp Lys Tyr Ile Val Ser Ala Ser Gly Asp Arg Thr
    370                 375                 380

Ile Lys Val Trp Ser Thr Ser Thr Cys Glu Phe Val Arg Thr Leu Asn
385                 390                 395                 400

Gly His Lys Arg Gly Ile Ala Cys Leu Gln Tyr Arg Asp Arg Leu Val
                405                 410                 415

Val Ser Gly Ser Ser Asp Asn Thr Ile Arg Leu Trp Asp Ile Glu Cys
            420                 425                 430

Gly Ala Cys Leu Arg Val Leu Glu Gly His Glu Glu Leu Val Arg Cys
        435                 440                 445

Ile Arg Phe Asp Asn Lys Arg Ile Val Ser Gly Ala Tyr Asp Gly Lys
    450                 455                 460

Ile Lys Val Trp Asp Leu Gln Ala Ala Leu Asp Pro Arg Ala Pro Ala
465                 470                 475                 480

Ser Thr Leu Cys Leu Arg Thr Leu Val Glu His Ser Gly Arg Val Phe
                485                 490                 495

Arg Leu Gln Phe Asp Glu Phe Gln Ile Ile Ser Ser Ser His Asp Asp
```

```
          500                 505                 510

Thr Ile Leu Ile Trp Asp Phe Leu Asn Val Pro Pro Ser Ala Gln Asn
        515                 520                 525

Glu Thr Arg Ser Pro Ser Arg Thr Tyr Thr Tyr Ile Ser Arg
    530                 535                 540


<210> 17
<211> 2151
<212> DNA
<213> Homo sapiens

<400> 17
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ggcacttcca gtatgattgt gcccaagcaa cggaaactct cagcaagcta tgaaaaggaa 360
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<210> 18
<211> 569
<212> PRT
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<213> Homo sapiens

<400> 18

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Asn Ser Ser Glu Arg Glu Asp Cys Asn Asn Gly Glu Pro Pro Arg Lys
            20                  25                  30

Ile Ile Pro Glu Lys Asn Ser Leu Arg Gln Thr Tyr Asn Ser Cys Ala
        35                  40                  45

Arg Leu Cys Leu Asn Gln Glu Thr Val Cys Leu Ala Ser Thr Ala Met
    50                  55                  60

Lys Thr Glu Asn Cys Val Ala Lys Thr Lys Leu Ala Asn Gly Thr Ser
65                  70                  75                  80

Ser Met Ile Val Pro Lys Gln Arg Lys Leu Ser Ala Ser Tyr Glu Lys
                85                  90                  95

Glu Lys Glu Leu Cys Val Lys Tyr Phe Glu Gln Trp Ser Glu Ser Asp
            100                 105                 110

Gln Val Glu Phe Val Glu His Leu Ile Ser Gln Met Cys His Tyr Gln
        115                 120                 125

His Gly His Ile Asn Ser Tyr Leu Lys Pro Met Leu Gln Arg Asp Phe
    130                 135                 140

Ile Thr Ala Leu Pro Ala Arg Gly Leu Asp His Ile Ala Glu Asn Ile
145                 150                 155                 160

Leu Ser Tyr Leu Asp Ala Lys Ser Leu Cys Ala Ala Glu Leu Val Cys
                165                 170                 175

Lys Glu Trp Tyr Arg Val Thr Ser Asp Gly Met Leu Trp Lys Lys Leu
            180                 185                 190

Ile Glu Arg Met Val Arg Thr Asp Ser Leu Trp Arg Gly Leu Ala Glu
        195                 200                 205

Arg Arg Gly Trp Gly Gln Tyr Leu Phe Lys Asn Lys Pro Pro Asp Gly
    210                 215                 220

Asn Ala Pro Pro Asn Ser Phe Tyr Arg Ala Leu Tyr Pro Lys Ile Ile
225                 230                 235                 240

Gln Asp Ile Glu Thr Ile Glu Ser Asn Trp Arg Cys Gly Arg His Ser
                245                 250                 255

Leu Gln Arg Ile His Cys Arg Ser Glu Thr Ser Lys Gly Val Tyr Cys
            260                 265                 270

Leu Gln Tyr Asp Asp Gln Lys Ile Val Ser Gly Leu Arg Asp Asn Thr
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```
        275                 280                 285

Ile Lys Ile Trp Asp Lys Asn Thr Leu Glu Cys Lys Arg Ile Leu Thr
    290                 295                 300

Gly His Thr Gly Ser Val Leu Cys Leu Gln Tyr Asp Glu Arg Val Ile
305                 310                 315                 320

Ile Thr Gly Ser Ser Asp Ser Thr Val Arg Val Trp Asp Val Asn Thr
                325                 330                 335

Gly Glu Met Leu Asn Thr Leu Ile His His Cys Glu Ala Val Leu His
            340                 345                 350

Leu Arg Phe Asn Asn Gly Met Met Val Thr Cys Ser Lys Asp Arg Ser
        355                 360                 365

Ile Ala Val Trp Asp Met Ala Ser Pro Thr Asp Ile Thr Leu Arg Arg
    370                 375                 380

Val Leu Val Gly His Arg Ala Ala Val Asn Val Val Asp Phe Asp Asp
385                 390                 395                 400

Lys Tyr Ile Val Ser Ala Ser Gly Asp Arg Thr Ile Lys Val Trp Asn
                405                 410                 415

Thr Ser Thr Cys Glu Phe Val Arg Thr Leu Asn Gly His Lys Arg Gly
            420                 425                 430

Ile Ala Cys Leu Gln Tyr Arg Asp Arg Leu Val Val Ser Gly Ser Ser
        435                 440                 445

Asp Asn Thr Ile Arg Leu Trp Asp Ile Glu Cys Gly Ala Cys Leu Arg
    450                 455                 460

Val Leu Glu Gly His Glu Glu Leu Val Arg Cys Ile Arg Phe Asp Asn
465                 470                 475                 480

Lys Arg Ile Val Ser Gly Ala Tyr Asp Gly Lys Ile Lys Val Trp Asp
                485                 490                 495

Leu Val Ala Ala Leu Asp Pro Arg Ala Pro Ala Gly Thr Leu Cys Leu
            500                 505                 510

Arg Thr Leu Val Glu His Ser Gly Arg Val Phe Arg Leu Gln Phe Asp
        515                 520                 525

Glu Phe Gln Ile Val Ser Ser Ser His Asp Asp Thr Ile Leu Ile Trp
    530                 535                 540

Asp Phe Leu Asn Asp Pro Ala Ala Gln Ala Glu Pro Pro Arg Ser Pro
545                 550                 555                 560

Ser Arg Thr Tyr Thr Tyr Ile Ser Arg
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<210> 19
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<212> PRT
<213> Homo sapiens

<400> 19
Asp Ser Gly Leu Asp Ser
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<210> 20
<211> 20
<212> PRT
<213> Homo sapiens

<400> 20
Ala Ala Val Asn Val Val Asp Phe Asp Asp Lys Tyr Ile Val Ser Ala
1 5 10 15

Ser Gly Asp Arg
20

<210> 21
<211> 17
<212> PRT
<213> Homo sapiens

<400> 21
Glu Leu Phe Pro Leu Ile Phe Pro Ala Glu Pro Ala Gln Ala Ser Gly
1 5 10 15

Pro

<210> 22
<211> 10
<212> PRT
<213> Homo sapiens

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<220>
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<222> (10)
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<400> 22
Cys Asp Arg His Asp Ser Gly Leu Asp Ser

1 5 10

<210> 23
<211> 4
<212> PRT
<213> Homo sapiens

<400> 23
Val Val Asn Val
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<210> 24
<211> 17
<212> PRT
<213> Homo sapiens

<400> 24
Ala Ala Val Asn Val Val Asp Phe Asp Asp Lys Tyr Ile Val Ser Ala
1 5 10 15
Ser

<210> 25
<211> 9
<212> PRT
<213> Homo sapiens

<400> 25
Leu Glu Gly His Glu Glu Leu Val Arg
1 5

<210> 26
<211> 12
<212> PRT
<213> Homo sapiens

<400> 26
Leu Val Val Ser Gly Ser Ser Asp Asn Thr Ile Arg
1 5 10

<210> 27
<211> 12
<212> PRT
<213> Homo sapiens

<400> 27
Ile Gln Asp Ile Glu Thr Ile Glu Ser Asn Trp Arg
1 5 10

<210> 28
<211> 9
<212> PRT
<213> Homo sapiens

<400> 28
Val Ile Ser Glu Gly Met Leu Trp Lys
1 5

<210> 29
<211> 10
<212> PRT
<213> Homo sapiens

<220>
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<222> (5)
<223> PHOSPHORYLATION

<220>
<221> MOD_RES
<222> (9)
<223> PHOSPHORYLATION

<400> 29
Asp Arg His Asp Ser Gly Leu Asp Ser Met
1 5 10

<210> 30
<211> 6
<212> PRT
<213> Homo sapiens

<220>
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<222> (2)
<223> PHOSPHORYLATION

<220>
<221> MOD_RES
<222> (6)
<223> PHOSPHORYLATION

<400> 30
Asp Ser Gly Leu Asp Ser
1 5